

SEOUENCE LISTING

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TECH CENTER 1600/2900

(1) GENERAL INFORMATION:

- (i) APPLICANT:
 - (A) NAME: Ferring B.V.
 - (B) STREET: Polaris Avenue 144
 - (C) CITY: PO Box 3129
 - (D) STATE: Hoofddorp
 - (E) COUNTRY: THE NETHERLANDS
 - (F) POSTAL CODE (ZIP): 2130 KC
- (ii) TITLE OF INVENTION: Agent for the Treatment of Osteoperosis
- (iii) NUMBER OF SEQUENCES: 7
- (iv) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
 - (v) CURRENT APPLICATION DATA:
 APPLICATION NUMBER: WO 99/****
- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

CTGCAGCTGC CTGAAGGAG

19

- (2) INFORMATION FOR SEQ ID NO: 2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

GGGCGGGCG GGGCTCTCG

19

- (2) INFORMATION FOR SEQ ID NO: 3:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
ATTCTACTGA CTTGGTGCGT G	21
(2) INFORMATION FOR SEQ ID NO: 4:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	0.1
GGAATATGTG CAACTTGGTG T	21
(2) INFORMATION FOR SEQ ID NO: 5:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: peptide	
<pre>(ix) FEATURE: (A) NAME/KEY: Modified-site (B) LOCATION:1 (D) OTHER INFORMATION:/product= "Glu in first position is pyroGLU"</pre>	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
Glu His Trp Ser Tyr Gly Leu Arg Pro Gly 1 5 10	
(2) INFORMATION FOR SEQ ID NO: 6:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: peptide	
<pre>(ix) FEATURE: (A) NAME/KEY: Modified-site (B) LOCATION:1 (D) OTHER INFORMATION:/product= "Glu in first position is pyroGlu"</pre>	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

Glu His Trp Ser His Gly Trp Tyr Pro Gly
1 5 10

- (2) INFORMATION FOR SEQ ID NO: 7:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 10 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION:1
 - (D) OTHER INFORMATION:/product= "Glu is pyroGlu"
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION:5
 - (D) OTHER INFORMATION:/product= "Xaa is His or Tyr"
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION: 7
 - (D) OTHER INFORMATION:/product= "Xaa is Trp or Leu"
 - (ix) FEATURE:
 - (A) NAME/KEY: Modified-site
 - (B) LOCATION:8
 - (D) OTHER INFORMATION:/product= "Xaa is Tyr or Arg"
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

Glu His Trp Ser Xaa Gly Xaa Xaa Pro Gly 1 5 10